

AMENDMENTS TO THE CLAIMS

The following listing of claims replaces all prior versions of claims in the application.

1. (Currently Amended) A wound dressing for accelerating epidermal regeneration which comprises:

at least one polypeptide (P) ~~having at least one species of epidermal regeneration-
accelerating minimal amino acid sequences (X) selected from the group consisting of Arg-Gly-
Asp (SEQ ID NO: 1), Ile-Lys-Val-Ala-Val (SEQ ID NO: 2), and Tyr-Ile-Gly-Ser-Arg (SEQ ID
NO: 3), and at least one auxiliary amino acid sequence (Y),~~

a polyalkylenepolyamine and/or polyarylenepolyamine (A) having a weight average
molecular weight of 2,000 to 60,000, and

a sheet (S) being polyurethane,

wherein the at least one polypeptide (P) is selected from the group consisting of:

(1) a polypeptide having 13 Arg Gly Asp sequences (SEQ ID NO: 1) and 13 (Gly
Ala Gly Ala Gly Ser)₉ sequences ((residues 1-6 of SEQ ID NO: 7)₉) chemically bonded
to each other in an alternating fashion,

(2) a polypeptide having 5 Arg Gly Asp sequences (SEQ ID NO: 1) and 5 (Gly
Ala Gly Ala Gly Ser)₃ sequences ((residues 1-6 of SEQ ID NO: 7)₃) chemically bonded to
each other in an alternating fashion,

(3) a polypeptide having 3 Arg Gly Asp sequences (SEQ ID NO: 1) and 3 (Gly
Val Pro Gly Val)₂ Gly Gly (Gly Ala Gly Ala Gly Ser)₃ sequences((residues 1-30 of SEQ
ID NO: 49)₉ chemically bonded to each other in an alternating fashion,

(4) a polypeptide having 13 Ile-Lys-Val-Ala-Val sequences (SEQ ID NO: 2) and 13 (Gly Ala Gly Ala Gly Ser)₉ sequences ((residues 1-6 of SEQ ID NO: 7)₉) chemically bonded to each other in an alternating fashion,

(5) a polypeptide having 5 Ile-Lys-Val-Ala-Val sequences (SEQ ID NO: 2) and 5 (Gly Ala Gly Ala Gly Ser)₃ sequences ((residues 1-6 of SEQ ID NO: 7)₃) chemically bonded to each other in an alternating fashion,

(6) a polypeptide having 3 Ile-Lys-Val-Ala-Val sequences (SEQ ID NO: 2) and 3 (Gly Val Pro Gly Val)₂ Gly Gly (Gly Ala Gly Ala Gly Ser)₃ sequences((residues 1-30 of SEQ ID NO: 49)₉ chemically bonded to each other in an alternating fashion,

(7) a polypeptide having 13 Tyr-Ile-Gly-Ser-Arg sequences (SEQ ID NO: 3) and 13 (Gly Ala Gly Ala Gly Ser)₉ sequences ((residues 1-6 of SEQ ID NO: 7)₉) chemically bonded to each other in an alternating fashion,

(8) a polypeptide having 5 Tyr-Ile-Gly-Ser-Arg sequences (SEQ ID NO: 3) and 5 (Gly Ala Gly Ala Gly Ser)₃ sequences ((residues 1-6 of SEQ ID NO: 7)₃) chemically bonded to each other in an alternating fashion, and

(9) a polypeptide having 3 Tyr-Ile-Gly-Ser-Arg sequences (SEQ ID NO: 3) and 3 (Gly Val Pro Gly Val)₂ Gly Gly (Gly Ala Gly Ala Gly Ser)₃ sequences((residues 1-30 of SEQ ID NO: 49)₉ chemically bonded to each other in an alternating fashion,

wherein the at least one polypeptide (P) and the sheet (S) are bonded by a covalent bonding,~~and~~

~~wherein said auxiliary amino acid sequence (Y) is selected from the group consisting of:~~

~~(Gly Ala)_a((residues 1-2 of SEQ ID NO: 4)_a),~~
~~(Gly Ala Gly Ala Gly Ser)_b((residues 1-6 of SEQ ID NO: 7)_b),~~
~~(Gly Ala Gly Ala Gly Tyr)_e((residues 1-6 of SEQ ID NO: 10)_e),~~
~~(Gly Ala Gly Val Gly Tyr)_d((residues 1-6 of SEQ ID NO: 13)_d),~~
~~(Gly Ala Gly Tyr Gly Val)_e((residues 1-6 of SEQ ID NO: 16)_e),~~
~~{Asp Gly Gly (Ala)_f Gly Gly Ala}_g((residues 1-12 of SEQ ID NO: 19)_g),~~
~~(Gly Val Pro Gly Val)_h((residues 1-5 of SEQ ID NO: 22)_h),~~
~~(Gly)_i((residue 1 of SEQ ID NO: 25)_i),~~
~~(Ala)_j((residue 1 of SEQ ID NO: 28)_j),~~
~~(Gly Gly Ala)_k((residues 1-3 of SEQ ID NO: 31)_k),~~
~~(Gly Val Gly Val Pro)_m((residues 1-5 of SEQ ID NO: 34)_m),~~
~~(Gly Pro Pro)_n((residues 1-3 of SEQ ID NO: 37)_n),~~
~~(Gly Ala Gln Gly Pro Ala Gly Pro Gly)_o((residues 1-9 of SEQ ID NO: 40)_o),~~
~~(Gly Ala Pro Gly Ala Pro Gly Ser Gln Gly Ala Pro Gly Leu Gln)_p((residues 1-15 of SEQ ID NO: 43)_p), and~~
~~(Gly Ala Pro Gly Thr Pro Gly Pro Gln Gly Leu Pro Gly Ser Pro)_q((residues 1-15 of SEQ ID NO: 46)_q),~~

~~wherein a is an integer from 5 to 100; b, c, d, and e each are an integer from 2 to 33; f is an integer from 1 to 194; g is an integer from 1 to {200/(6 + f)} with any fraction omitted; h is an integer from 2 to 40; i and j each are an integer from 10 to 200; k is an~~

~~integer from 3 to 66; m is an integer from 2 to 40; n is an integer from 3 to 66; o is an integer from 1 to 22; and p and q each are an integer from 1 to 13.~~

2-6. (Cancelled)

7. (Original) The wound dressing according to Claim 1

wherein the polyalkylenepolyamine and/or polyarylenepolyamine (A) is a polyethyleneimine.

8. (Withdrawn) A method for epidermal regeneration treatment which comprises using the wound dressing according to Claim 1.

9-13. (Cancelled)

14. (Currently Amended) The wound dressing according to claim 1 [[13]], wherein the at least one polypeptide (P) is selected from the group consisting of ~~ProNectin F and ProNectin L~~

the polypeptide having 13 Arg Gly Asp sequences (SEQ ID NO: 1) and 13 (Gly Ala Gly Ala Gly Ser)₉ sequences ((residues 1-6 of SEQ ID NO: 7)₉) chemically bonded to each other in an alternating fashion, and

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the polypeptide having 13 Ile-Lys-Val-Ala-Val sequences (SEQ ID NO: 2) and 13 (Gly Ala Gly Ala Gly Ser)₉ sequences ((residues 1-6 of SEQ ID NO: 7)₉) chemically bonded to each other in an alternating fashion.